



Docket No.: 20347 US1 (C38435/128985)

IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

In re Application of :)
Heinrich BACHMANN, *et al.*)
Serial No.: 10/053,192) Examiner: not yet assigned
Filed: January 15, 2002) Art Unit: 1652
For: **β,β -CAROTENE 15,15'-**)
MONOOXYGENASES, NUCLEIC)
ACID SEQUENCES CODING)
THEREFOR AND THEIR USE)

New York, NY
September 30, 2002

TRANSMITTAL OF PROPOSED DRAWING CORRECTION

Box Missing Parts
Commissioner For Patents
Washington, D.C. 20231
Sir:

In accordance with 37 CFR § 121(d) and MPEP § 608.02(q) and at the request of the Examiner, (see Notice to Filing Missing Parts mailed April 29, 2002), the attached corrected drawing figures (see Exhibit 1) are submitted for consideration. It is respectfully requested that the Examiner enter corrected figures 2-8 in this application.

The correction to figures 2-5 is as follows:

- a. figures 2-5 have been reformatted to correct the sizes of the margins in order to conform to the requirements of 37 CFR § 1.84(g).

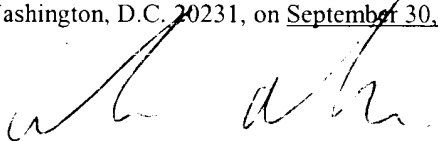
The corrections to figures 6-8 are as follows:

- a. figures 7 and 8 have been reformatted to correct the sizes of the margins in order to conform to the requirements of 37 CFR § 1.84(g).
- b. all descriptive text, other than text indicating the figure number, has been cancelled from figures 6-8 in order to conform to the requirements of 37 CFR § 1.84(o).

For the convenience of the Examiner, we attach as Exhibit 2 a copy of figures 6-8 as amended (without the redline).

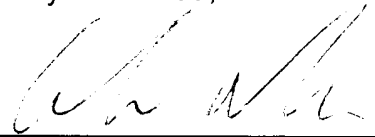
It is submitted that the entry of the corrections to Figures 2-8 does not introduce new matter, and is proper. See 37 CFR § 121(a)(6). Accordingly, entry of the attached corrected drawing figures in the above referenced application, respectfully is requested. If the Examiner has any questions regarding this paper, please contact the undersigned.

I hereby certify that this correspondence is being deposited with the United States Postal Service with sufficient postage as first class mail in an envelope addressed to Box Missing Parts, Commissioner For Patents, Washington, D.C. 20231, on September 30, 2002.



Gonzalo Merino

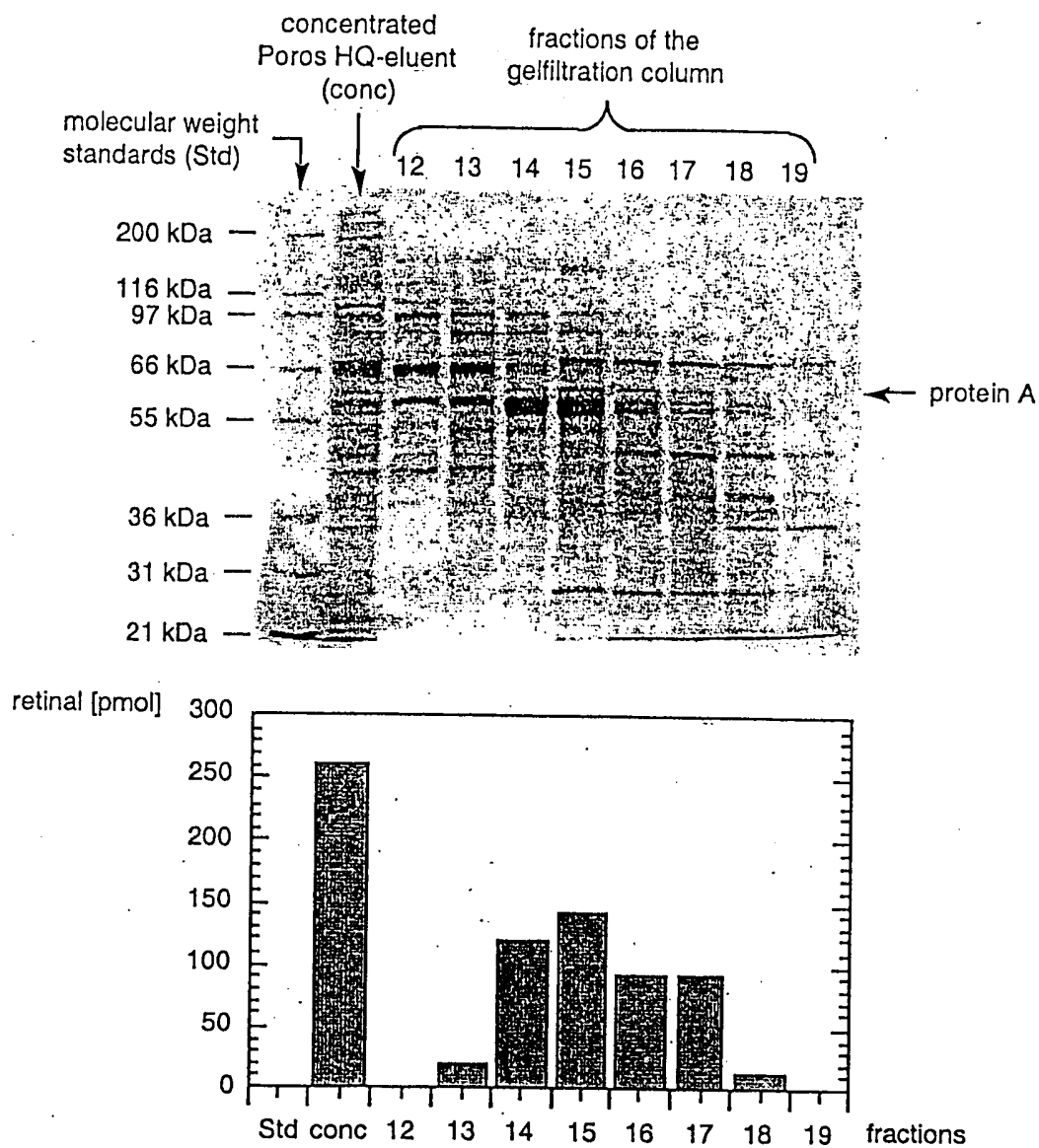
Respectfully submitted,

By: 

Gonzalo Merino
Registration No. 51,192
BRYAN CAVE LLP
245 Park Avenue
New York, NY 10167-0034
(212) 692-1800



Figure 1



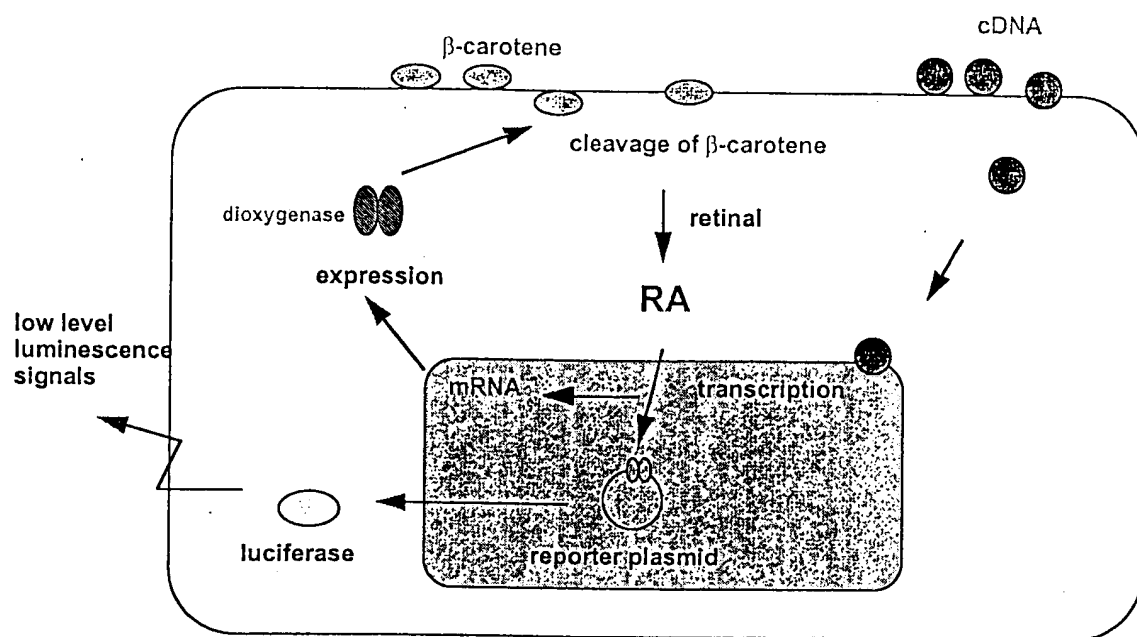


Figure 2



Figure 3
No. 2

Seq. ID

1 CGGATCCACT AGTAACGGCC GCCAGTGTGG TGGAATCCAT
CCTTCTATGT

51 AACAGGAAAG AGCTGTTCTT AGCCCAGAGA GGAGGGCACC
GTACGCCTGC

101 AGGAGCAGCT GGGTAGAGGA CACAGGAGAG CGATGGAGAC
AATATTTAAC

151 AGAAACAAAG AAGAGCATCC AGAGCCCATA AAAGCTGAGG
TGCAAGGTCA

201 GTTGCCCACT TGGTTGCAAG GGGTACTTCT CCGAAATGGC
CCAGGGATGC

251 ACACAATAGG GGACACTAAA TACAACCACT GGTTTGATGG
CTTGGCTCTG

301 CTGCACAGCT TCACGTTTAA AAATGGTGAA GTTTACTACA
GAAGTAAGTA

351 CCTCCGAAGT GACACATACA ACTGCAATAT AGAAGCAAAC
CGAATCGTGG

401 TGTCTGAGTT TGGAACCATG GCTTATCCGG ATCCATGCAA
AAACATATTT

451 GCCAAGGCAT TCTCATACTT ATCTCACACC ATTCCTGAGT
TCACGGACAA

501 CTGCCTGATC AACATTATGA AAAGTGGGGA TGATTATTAT
GCTACCACTG

551 AGACTAACTT CATCAGAAAA ATTGATCCAC AGACTCTGGA
GACTAGAT

601 AAGGTAGACT ACAGCAAATA TGTAGCTGTA AACTTGGCAA
CTTCTCACCC

651 ACACTATGAC AGTGCTGGAA ATATTCTCAA CATGGGTACT
TCAATTGTTG

701 ATAAAGGGAG AACAAAATAT GTTCTCTTTA AGATCCCTTC
CTCTGTACCA

751 GAAAAAGAAA AGAAGAAATC TTGTTTTAAA CACCTGGAAG
TAGTATGCTC

801 CATCCCTTCT CGCTCCCTGC TCCAACCAAG CTACTACCAC
AGCTTTGGAA

851 TCACAGAAAA TTATATTGTG TTCATAGAGC AGCCATTTAA
ACTGGATATT



901 GTCAAAGTGG CAACTGCCTA CATCCGAGGT GTGAACTGGG
CTTCCTGCCT

951 TTCCTTTCAT AAGGAGGATA AGACGTGGTT TCACTTTGTA
GACAGAAAGA

1001 CGAAAAAAGA AGTATCCACC AAGTTTTACA CTGATGCTTT
GGTGCTTTAT

1051 CACCACATAA ATGCTTACGA AGAAGATGGC CACGTTGTTT
TTGATATCGT

1101 TGCCTACAGA GACAATAGCT TGTACGATAT GTTTTACTTA
AAAAAACTGG

1151 ACAAAGACTT TGAAGTGAAC AACAAGCTTA CCTCCATCCC
AACCTGCAAG

1201 CGCTTTGTTG TGCCTCTGCA GTATGACAAG GATGCAGAAG
TAGGTTCTAA

1251 TTTAGTCAAA CTTCCAACCT CCGCAACTGC TGTAAAAGAA
AAAGATGGCA

1301 GCATCTATTG TCAACCTGAA ATATTATGTG AAGGGATAGA
ACTGCCTCGT

1351 GTCAACTATG ACTACAATGG CAAAAAATAC AAGTATGTCT
ATGCAACAGA

1401 AGTCCAGTGG AGCCCAGTTC CTACAAAGAT TGCAAACTG
AATGTCCAAA

1451 CAAAGGAAGT ACTGCACTGG GGAGAAGACC ACTGCTGGCC
CTCAGAGCCC

1501 ATCTTTGTTC CCAGCCCCGA TGCAAGAGAA GAGGATGAAG
GTGTTGTTTT

1551 GACCTGTGTT GTGGTGTCTG AGCCAAATAA AGCACCTTC
CTACTCATCT

1601 TGGATGCTAA AACATTCAAA GAATTGGGCC GAGCCACAGT
TAACGTAGAA

1651 ATGCATCTGG ACCTGCATGG GATGTTTATA CCACAGAATG
ATTTGGGGGC

1701 TGAGACGGAA TAAAACGCTA TTGATCCGAC TACACAACT
GAGACAACTT

1751 TCTACTGAAC ATGAGTTAAT ATCCCTTTTA CCATTCAAGA
ACAACCATAT

1801 AACGACACAA AATGACTATG TATAATCTCT TAAATAATAG
ATATAATCCT

1851 TTTAAGGCAC AGCGATGAGT TTTACTACAG GTAACGATAT
GCACAACTGG



1901 CATATAACTA TTCCAAAAGA AGAAGAACGA TCAGTGTTTT
AGAAGTGCTA

1951 ATGTTGTACA TAACGGCGGC AGAGGGAACA GGAGAGAAAG
GTAACGGGAA

2001 TATTTAATAG AATATAGATT TCTGAGCAAA TGAAGTGCAG
TATTTATGGT

2051 GTGATGCATG GCATGAGTCA CATAGGTCTG CAGCTCATGT
ATCTTTTAGA

2101 GATCGTTTCA AGATTGCAGC TTGTGATGCA AGTTTTCTCC
AGCCAGAAAA

2151 CCTCATTTTA AACCATCTGC TACTGGTAAT TCATACCAAT
GCATTTTCTT

2201 GGTGCTCGAT TTACACTATA ACCAAAGTTA AGTATTACAT
TCAGGTGCTA

2251 CAACTTTCTA ATTTACAACC GAAACAAACA AGCAAACAGC
ACTTGCTTTG

2301 CTAATAACCC CATGGTGTAT TTTTCCTTTT TATGATGACA
AAACCAAGTA

2351 CATATGGTTT TATGTAGCAT TCAATTATAC TTCAGTGCTA
TTCCATCCTA

2401 ATGTTATAAG CAATTTGTAT TTAAATCAGT TTCCTTGAG
AATATCTGAC

2451 ATAACATTTT GTGTAATGAG ATGACTATGT TGTCTAAAGA
TGAACAGGAA

2501 TGTATCTTTT ATTAGTATTG TTAATTGTGT TACTAATACT
ATGCATATGA

2551 ATGAGAGCAA TGTATTTCTA GGAGAACTCA GATATACATT
CAACAATTTT

2601 TGTAGGTGAA AATGCATTTA CTGATGAAAG TTGAATCGTT
AATGAGGGAG

2651 AAAACTGGGT ATCCATCCAT CCAACTATGT TAGGTGTTCA
CCTGGTCTGT

2701 ATGTGACACC ACGCTGTTTG GGTATCTCTC ACTTTCACAT
ACCTGTTCTC

2751 ATGGTTTCTG CTACTCACTG TATTTTGCAG GAGAGAAACA
AAATGAAATC

2801 ACTGTCACCT ACTATCGCCC CATCACATAA GAACAATGGG
GCTTTGGTGA



2851 CTTGTTTCATG ATTACATAAG ATGTTTGCAG CAGAGCAGCA
ATAGAACCAA

2901 CACCATCCAC AGTTCTTGCT TGCTCTGTTA TGACTCCCTT
TGCTGTCTTT

2951 ATGGTTTGCA TGTATGAAGA ATACACTGCC TAATTCTAAT
GTTAAAAAGT

3001 CACTGGGGTC AGATCTAGAG CTTAAGTAAG CAGTCTGGGG
TTTTCAAATG

3051 TTTATATGTT CCATAAAATG GAAATAAACA CCTCCATAAT
AAAAAAAAAA

3101 AAAAAAAAAA A



Figure 4
No. 1

Seq. ID

1 METIFNRNKE EHPEPIKAEV QGQLPTWLQG VLLRNGPGMH
TIGDTKYNHV

51 FDGLALLHSF TFKNGEVYR SKYLRSDTYN CNIEANRIVV
SEFGTMAYPD

101 PCKNIFAKAF SYLSHTIPEF TDNCLINIMK TGDDYYATSE
TNFIRKIDPQ

151 TLETLDKVDY SKYVAVNLAT SHPHYDSAGN ILNMGTSIVD
KGRTKYVLFK

201 IPSSVPEKEK KKSCFKHLEV VCSIPSRSL QPSYYHSFGI
TENYIVFIEQ

251 PFKLDIVKLA TAYIRGVNWA SCLSFHKEDK TWFHFVDRKT
KKEVSTKFYT

301 DALVLYHHIN AYEEDGHVVF DIVAYRDNLS YDMFYLLKLD
KDFEVNNKLT

351 SIPTCKRFVV PLQYDKDAEV GSNLVKLPTS ATAVKEKDGS
IYCQPEILCE

401 GIELPRVNYD YNGKKYKYVY ATEVQWSPVP TKIAKLVNQT
KEVLHWGEDH

451 CWPSEPIFVP SPDAREEDED VVLTCCVVSE PNKAPFLLIL
DAKTFKELGR

501 ATVNVEMHLD LHGMFIPQND LGAETE

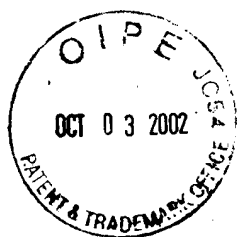


Figure 5

Seq ID No. 4 and Seq ID No. 5

10 EEHP EPIKA EVQGQLPTWLQGVLLR..NGPGMHTIGDTKYNHWF DGLALL
57
20 EELSSPLTAHV TGRIP LWTGSLLR CFTGPGLFEVGSEPFYHLFDGQALL
69
58 HSFTFKNGE VYYRSKYLRSDTYNCNIEANRIVVSEFG..TMAYPDPCKNI
105
70 HKFDFKEGHV TYHRRFIRTDAYVRAMTEKRIVITEFGFTTCAFPDPCKNI
119
106 FAKAFSYLSHTIPEFTDNCLINIMKTGDDYYATSETNFIRKIDPQTLETL
155
120 FSRFFSYFRGV..EVT DNALVNVYPVGEDYYACTETNFITKINPETLETI
167
156 ..DKVDYSKYVAVNLATSHPHYDSAGN ILNMGT SIVDKGR TKYVLFKIPS
203
168 FTKQVDLCNYVSVNGATAHPHIENDGT VYNIGNCFGKNFSIAYNIVKIPP
217
204 SVPEKEKKKSCFKHLEVVC SIPSRSL LQPSYYHSFGITENYIVFIEQPFK
253
218 LQADKEDPISKFTS.EIVVQFP CSDRFKPSYVHSFGLTPNYIVFVETPVK
266
254 LDIVKLATAY.IRGVNWASCL.SFHKEDK.TWFHFVDRKTKKEVSTKFYT
300
267 INLFKFLSSWSLWGANYMDCFESFTNETMGVWLHIADKKRKKYLNNKYRT
316
301 DALVLYHHINAYEEDGHVVF DIVAYRDN SL...YDMFY LKKLDKDFE...
344
317 SPFNLFHHINTYEDNGFLIVDLCCWKGF EFVYNYFTLYLANLRENWEEVK
366
345 VNNKLTSIPTCKRFV VPLQYDKDAEVGSNLVKLP.TSATAV..KEKD GSI
391
367 KNARKAPQPEVRRYVLPLNIDK.ADTGKNLVTL PNTTATAILCSDEFTTI
415
392 YCQPEILCEG....IELPRVNYD.YNGKKYKYVYATEVQWSPVPTKIAKL
436
416 WLEPEVLFSGPRQAFEFQIN YQKYCGKPYTYAYGLGLNHF.VPDR LCKL
464



437 NVQTKEVLH..WGEDHCWPSEPIFVSPDAREEDEGVVLTCTVVVSEPNKA
484 |.||| | | :||| ||| |||:||||. ||
465 NVKTKETWFTVWQEPDSYPSEPIFVSHPDAL EDDGVVLSVVVSPGAGQK
514
485 P.FLLILDAKTFKELGRA..TVNVEMHLDLHG MF 515
 | :|||.|| |. || || : . . ||:|
515 PAYLLILNAKDLSEVARAEFTVEINIPVTFHGLF 548

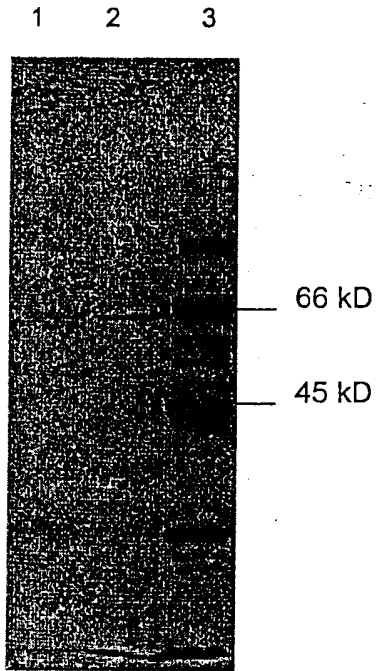


Fig. 6

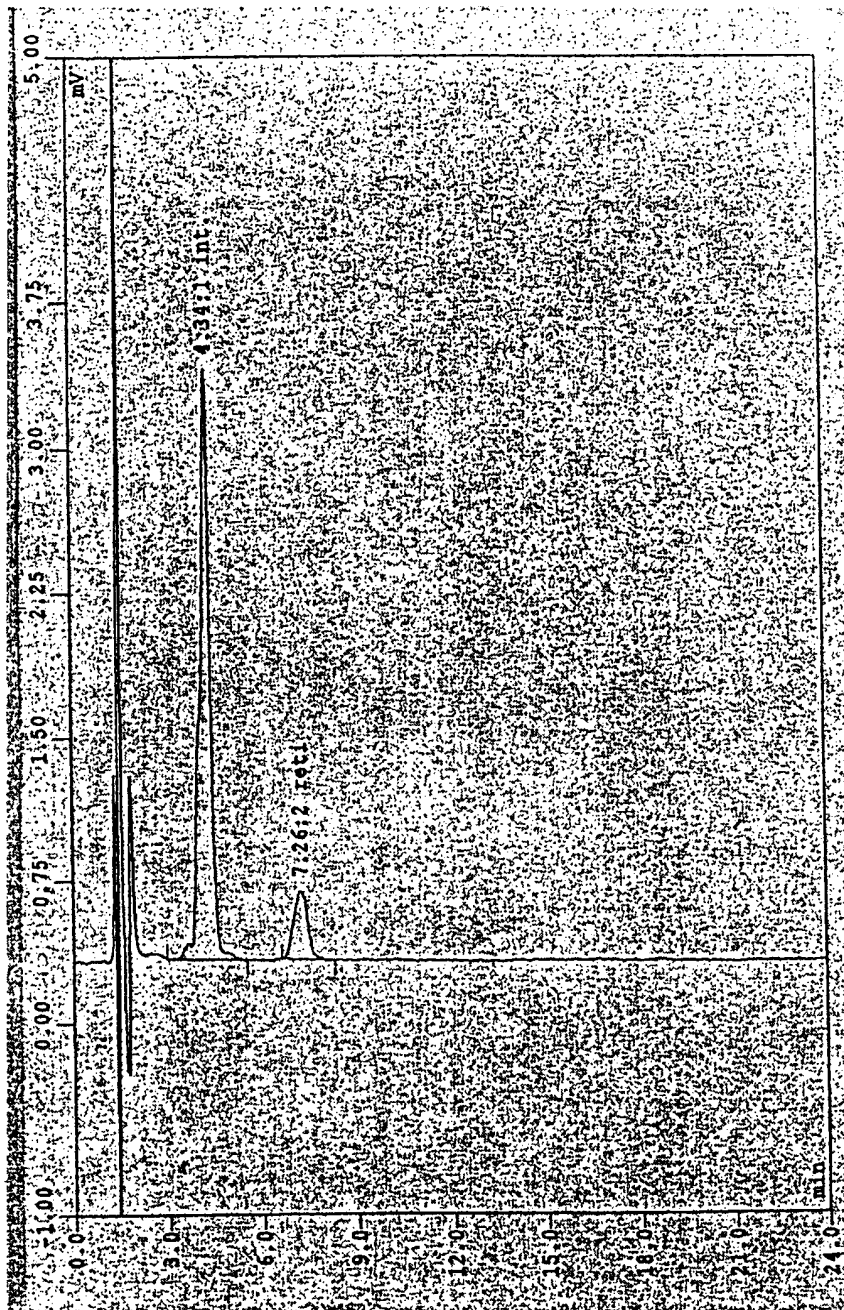


Fig. 7

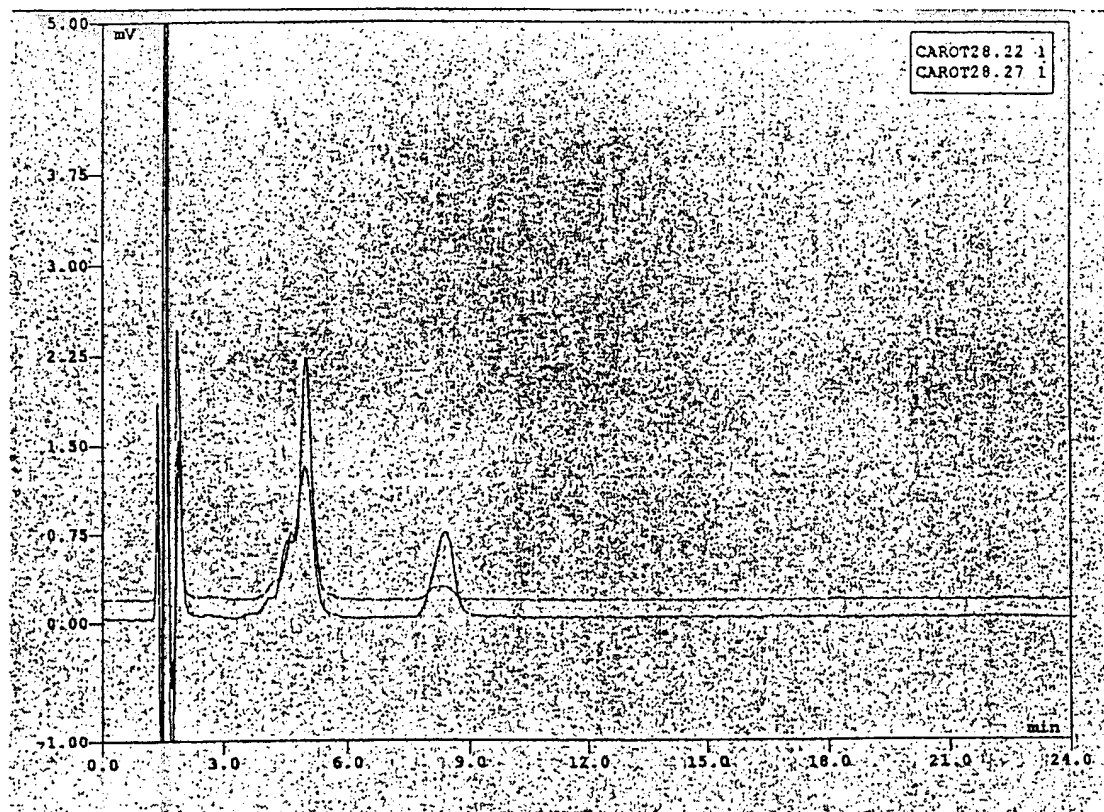
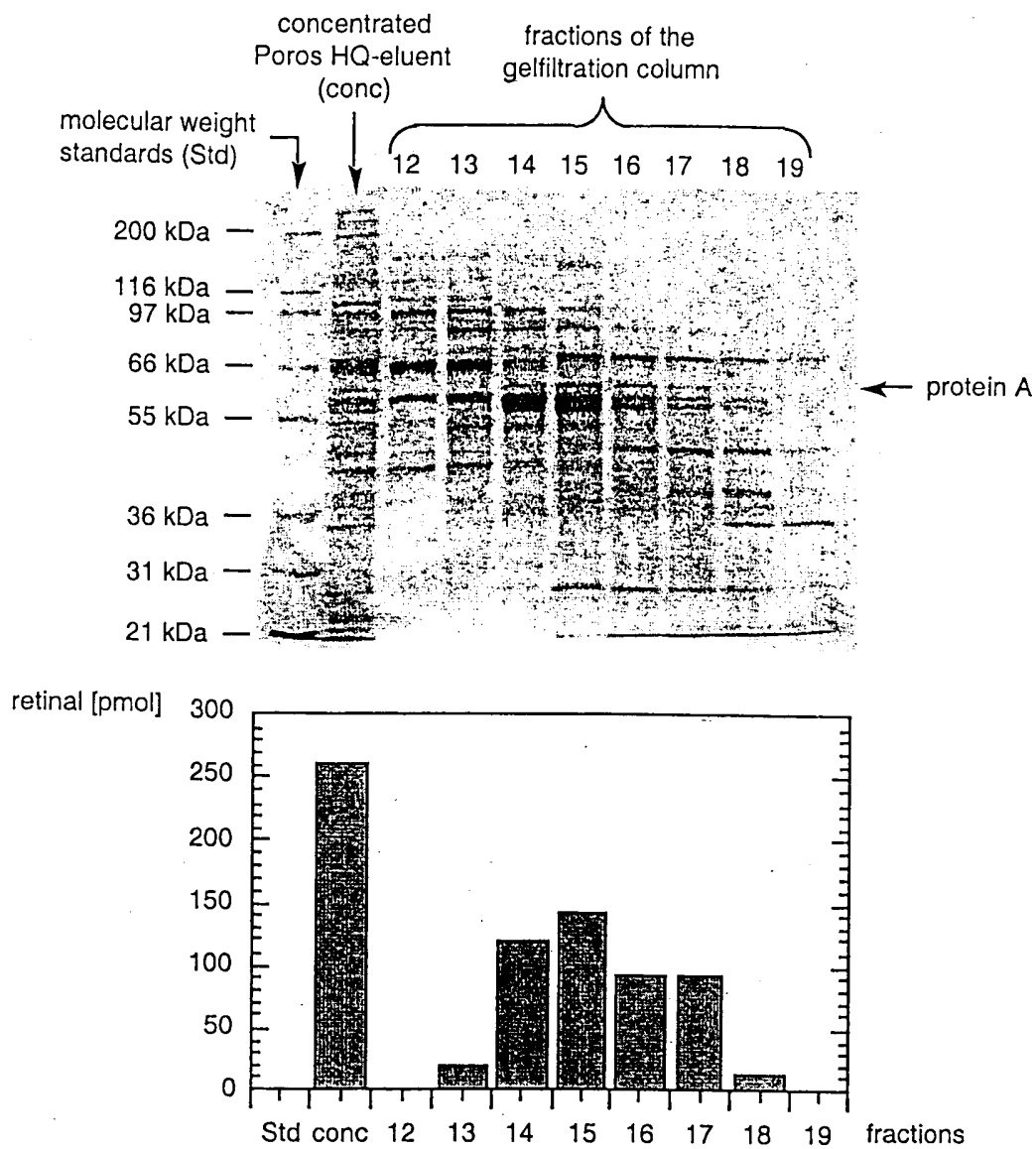


Fig. 8

Figure 1



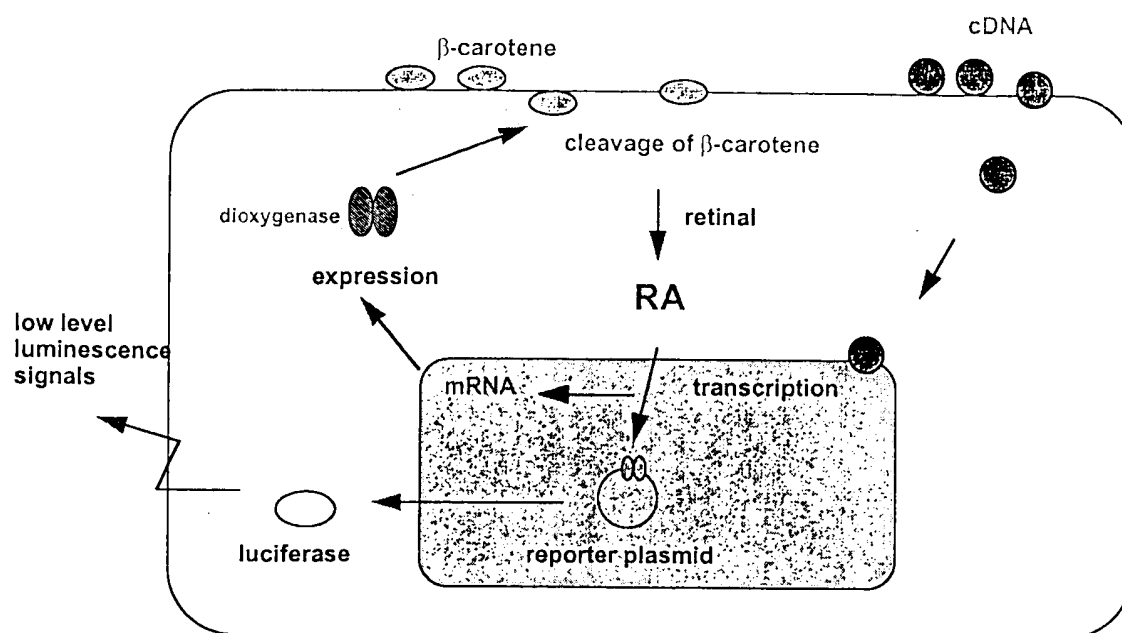


Figure 2



Figure 3
No. 2

Seq. ID

1 CGGATCCACT AGTAACGGCC GCCAGTGTGG TGAATCCAT
CCTTCTATGT

51 AACAGGAAAG AGCTGTTCTT AGCCCAGAGA GGAGGGCACC
GTACGCCTGC

101 AGGAGCAGCT GGGTAGAGGA CACAGGAGAG CGATGGAGAC
AATATTTAAC

151 AGAAACAAAG AAGAGCATCC AGAGCCCATA AAAGCTGAGG
TGCAAGGTCA

201 GTTGCCCACT TGGTTGCAAG GGGTACTTCT CCGAAATGGC
CCAGGGATGC

251 ACACAATAGG GGACACTAAA TACAACCACT GGTTTGATGG
CTTGGCTCTG

301 CTGCACAGCT TCACGTTTAA AAATGGTGAA GTTTACTACA
GAAGTAAGTA

351 CCTCCGAAGT GACACATACA ACTGCAATAT AGAAGCAAAC
CGAATCGTGG

401 TGTCTGAGTT TGGAACCATG GCTTATCCGG ATCCATGCAA
AAACATATTT

451 GCCAAGGCAT TCTCATACTT ATCTCACACC ATTCCTGAGT
TCACGGACAA

501 CTGCCTGATC AACATTATGA AAAGTGGGGA TGATTATTAT
GCTACCACTG

551 AGACTAACTT CATCAGAAAA ATTGATCCAC AGACTCTGGA
GACACTAGAT

601 AAGGTAGACT ACAGCAAATA TGTAAGCTGTA AACTTGGCAA
CTTCTCACCC

651 ACACTATGAC AGTGCTGGAA ATATTCTCAA CATGGGTACT
TCAATTGTTG

701 ATAAAGGGAG AACAAAATAT GTTCTCTTTA AGATCCCTTC
CTCTGTACCA

751 GAAAAAGAAA AGAAGAAATC TTGTTTTAAA CACCTGGAAG
TAGTATGCTC

801 CATCCCTTCT CGCTCCCTGC TCCAACCAAG CTACTACCAC
AGCTTTGGAA

851 TCACAGAAAA TTATATTGTG TTCATAGAGC AGCCATTTAA
ACTGGATATT



901 GTCAAAGTGG CAACTGCCTA CATCCGAGGT GTGAACTGGG
CTTCCTGCCT

951 TTCCTTTCAT AAGGAGGATA AGACGTGGTT TCACTTTGTA
GACAGAAAGA

1001 CGAAAAAAGA AGTATCCACC AAGTTTTACA CTGATGCTTT
GGTGCTTTAT

1051 CACCACATAA ATGCTTACGA AGAAGATGGC CACGTTGTTT
TTGATATCGT

1101 TGCCTACAGA GACAATAGCT TGTACGATAT GTTTTACTTA
AAAAAAGTGG

1151 ACAAAGACTT TGAAGTGAAC AACAAAGCTTA CCTCCATCCC
AACCTGCAAG

1201 CGCTTTGTTG TGCCTCTGCA GTATGACAAG GATGCAGAAG
TAGGTTCTAA

1251 TTTAGTCAAA CTTCCTCACTT CCGCAACTGC TGTAAGAGAA
AAAGATGGCA

1301 GCATCTATTG TCAACCTGAA ATATTATGTG AAGGGATAGA
ACTGCCTCGT

1351 GTCAAAGTATG ACTACAATGG CAAAAAATAC AAGTATGTCT
ATGCAACAGA

1401 AGTCCAGTGG AGCCAGTTC CTACAAAGAT TGCAAAAGTGG
AATGTCCAAA

1451 CAAAGGAAGT ACTGCACTGG GGAGAAGACC ACTGCTGGCC
CTCAGAGCCC

1501 ATCTTTGTTC CCAGCCCCGA TGCAAGAGAA GAGGATGAAG
GTGTTGTTTT

1551 GACCTGTGTT GTGGTGTCTG AGCCAAATAA AGCACCTTC
CTACTCATCT

1601 TGGATGCTAA AACATTCAAA GAATTGGGCC GAGCCACAGT
TAACGTAGAA

1651 ATGCATCTGG ACCTGCATGG GATGTTTATA CCACAGAATG
ATTGGGGGC

1701 TGAGACGGAA TAAAACGCTA TTGATCCGAC TACACAACT
GAGACAACTT

1751 TCTACTGAAC ATGAGTTAAT ATCCCTTTTA CCATTCAAGA
ACAACCATAT

1801 AACGACACAA AATGACTATG TATAATCTCT TAAATAATAG
ATATAATCCT

1851 TTTAAGGCAC AGCGATGAGT TTTACTACAG GTAACGATAT
GCACAACTGG



1901 CATATACTA TTCCAAAAGA AGAAGAACGA TCAGTGT TTTT
AGAAGTGCTA

1951 ATGTTGTACA TAACGGCGGC AGAGGGAACA GGAGAGAAAG
GTAACGGGAA

2001 TATTTAATAG AATATAGATT TCTGAGCAAA TGAAGTGCAG
TATTTATGGT

2051 GTGATGCATG GCATGAGTCA CATAGGTCTG CAGCTCATGT
ATCTTTTAGA

2101 GATCGTTTCA AGATTGCAGC TTGTGATGCA AGTTTTCTCC
AGCCAGAAAA

2151 CCTCATTTTA AACCATCTGC TACTGGTAAT TCATACCAAT
GCATTTTCTT

2201 GGTGCTCGAT TTACACTATA ACCAAAGTTA AGTATTACAT
TCAGGTGCTA

2251 CAACTTTCTA ATTTACAACC GAAACAAACA AGCAAACAGC
ACTTGCTTTG

2301 CTAATAACCC CATGGTGTAT TTTTCCTTTT TATGATGACA
AAACCAAGTA

2351 CATATGGTTT TATGTAGCAT TCAATTATAC TTCAGTGCTA
TTCCATCCTA

2401 ATGTTATAAG CAATTTGTAT TTAAATCAGT TTTCTTGAG
AATATCTGAC

2451 ATAACATTTT GTGTAATGAG ATGACTATGT TGTCTAAAGA
TGAACAGGAA

2501 TGTATCTTTT ATTAGTATTG TTAATTGTGT TACTAATACT
ATGCATATGA

2551 ATGAGAGCAA TGTATTTCTA GGAGAACTCA GATATACATT
CAACAATTTT

2601 TGTAGGTGAA AATGCATTTA CTGATGAAAG TTGAATCGTT
AATGAGGGAG

2651 AAAACTGGGT ATCCATCCAT CCAACTATGT TAGGTGTTCA
CCTGGTCTGT

2701 ATGTGACACC ACGCTGTTTG GGTATCTCTC ACTTTACAT
ACCTGTTCTC

2751 ATGGTTTCTG CTA CTACTACTG TATTTTGCAG GAGAGAAACA
AAATGAAATC

2801 ACTGTCACTT ACTATCGCCC CATCACATAA GAACAATGGG
GCTTTGGTGA



2851 CTTGTTTCATG ATTACATAAG ATGTTTGCAG CAGAGCAGCA
ATAGAACCAA

2901 CACCATCCAC AGTTCTTGCT TGCTCTGTGA TGA CTCCCTT
TGCTGTCTTT

2951 ATGGTTTGCA TGTATGAAGA ATACACTGCC TAATTCTAAT
GTTAAAAAGT

3001 CACTGGGGTC AGATCTAGAG CTTAAGTAAG CAGTCTGGGG
TTTCAAATG

3051 TTTATATGTT CCATAAAATG GAAATAAACA CCTCCATAAT
AAAAAAAAAA

3101 AAAAAAAAAA A



Figure 4
No. 1

Seq. ID

1 METIFNRNKE EHPEPIKAEV QGQLPTWLQG VLLRNGPGMH
TIGDTKYNHW

51 FDGLALLHSF TFKNGEVYR SKYLRSDTYN CNIEANRIVV
SEFGTMAYPD

101 PCKNIFAKAF SYLSHTIPEF TDNCLINIMK TGDDYYATSE
TNFIRKIDPQ

151 TLETLDKVDY SKYVAVNLAT SHPHYDSAGN ILNMGTSIVD
KGRTKYVLFK

201 IPSSVPEKEK KKSCFKHLEV VCSIPSRLL QPSYYHSFGI
TENYIVFIEQ

251 PFKLDIVKLA TAYIRGVNWA SCLSFHKEDK TWFHFVDRKT
KKEVSTKFYT

301 DALVLYHHIN AYEEDGHVVF DIVAYRDNLS YDMFYLKKLD
KDFEVNNKLT

351 SIPTCKRFVV PLQYDKDAEV GSNLVKLPTS ATAVKEKDGS
IYCQPEILCE

401 GIELPRVNYD YNGKKYKYVY ATEVQWSPVP TKIAKLVQQT
KEVLHWGEDH

451 CWPSEPIFVP SPDAREEDEG VVLTCVVVSE PNKAPFLIL
DAKTFKELGR

501 ATVNVEMHLD LHGMFIPQND LGAETE



Figure 5

Seq ID No. 4 and Seq ID No. 5

57 10 EEHPEPIKAĖVQQLPTWLQGVLLR..NGPGMHTIGDTKYNHWFDGLALL
|| | : | | | : | | | | | : | : | | | | |
69 20 EELSSPLTAHVGTGRIPWLWTGSLRLRCFTGPGLFEVGSEPFYHLFDGQALL
58 HSFTFKNGEVYYSKYLRSDTYNCNIEANRIVVSEFG..TMAYPDPCCKNI
105 | | | | | : : : | | . | | : : | | | | | : | | | | |
119 70 HKFDFKEGHVITYHRRFIRTDAYVRAMTEKRIVITEFGFTTCAFPDPCCKNI
120 FAKAFSYLSHTIPEFTDNCLINIMKTGDDYYATSETNFIRKIDPQTLETI
155 | : : | | | | | | : : | : | | | . | | | | | : | | : | | : |
167 120 FSRFFSYFRGV..EVDNALVNVPVGEDYYACTETNFITKINPETLETI
156 ..DKVDYSKYVAVNLATSHPHYDSAGNILNMGTSIVDKGRTKYVLFKIPS
203 . | | | | . | | | | : . | : | . | | : | |
217 168 FTKQVDLCNYVSVNGATAHPHIENDGTVYNIGNCFGKNFSIAYNIVKIPP
204 SVPEKEKKKSCFKHLEVVCŠIPSRLLQPSYYHSFGITENYIVFIEQPFK
253 : | | | | | : | | | . | | | | | : | | | | | : | | |
266 218 LQADKEDPISKFTS.EIVVQFPCSDRFKPSYVHSFGLTPNYIVFVETPVK
254 LDIVKLATAY.IRGVNWASCL.SFHKEDK.TWFHFVDRKTKKEVSTKFYT
300 : : | . : : | : | | | | | : | | | . : |
316 267 INLFKFLSSWSLWGANYMDCFESFTNETMGVWLHIADKKRKKYLNNKYRT
301 DALVLYHHINAYEEDGHVVFIDIVAYRDNŠL...YDMFYLKKLKDKE...
344 | : | | | | : : | : : | | | | | : : |
366 317 SPFNLFHHINTYEDNGFLIVDLCCWKGFEFVYNYFTLYLANLRENWEEVK
345 VNNKLTŠIPTCKRFVVPLQYDKDAEVGSNLVKLP.TSATAV..KEKDGSİ
391 | : | : : | | | : | | | | | . | | : : . |
415 367 KNARKAPQPEVRRYVLPLNIDK.ADTGKNLVTLPNTTATAILCSDEFTTI
392 YCQPEILCEG....IELPRVNYD.YNGKKYKYVYATEVQWSPVPTKIAKL
436 : : | : | | : : | | | | | . | | : : |
464 416 WLEPEVLFSGPRQAFEFQINQKYCGKPYTYAYGLGLNHF.VPDRLCKL



437 NVQTKEVLH..WGEDHCWPSEPIFVPSPDAREEDEGVVLTCTVVVSEPNKA
484 ||.||| || :||||||| ||| |||:|||||. ||
465 NVKTKETWFTVWQEPDSYPSEPIFVSHPDAL EDDGVVLSVVVSPGAGQK
514
485 P.FLLILDAKTFKELGRA..TVNVEMHLDLHG MF 515
 |:|||||.|| |. || ||: . ||:|
515 PAYLLILNAKDLSEVARAEFTVEINIPVTFHGLF 548

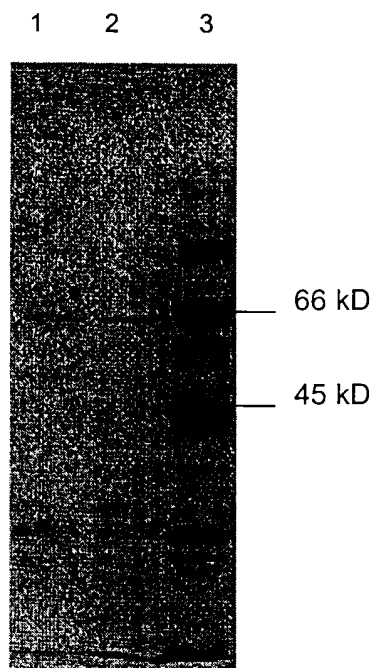


Fig. 6 shows a 10% polyacrylamide gel with E.coli expressed β,β -carotene 15,15'-monooxygenase after affinity tag purification; lane 1 and lane 2: 2 fractions from the Co^{2+} -chelate column showing the main band at 60 kD; lane 3: low range molecular weight marker (Bio Rad).

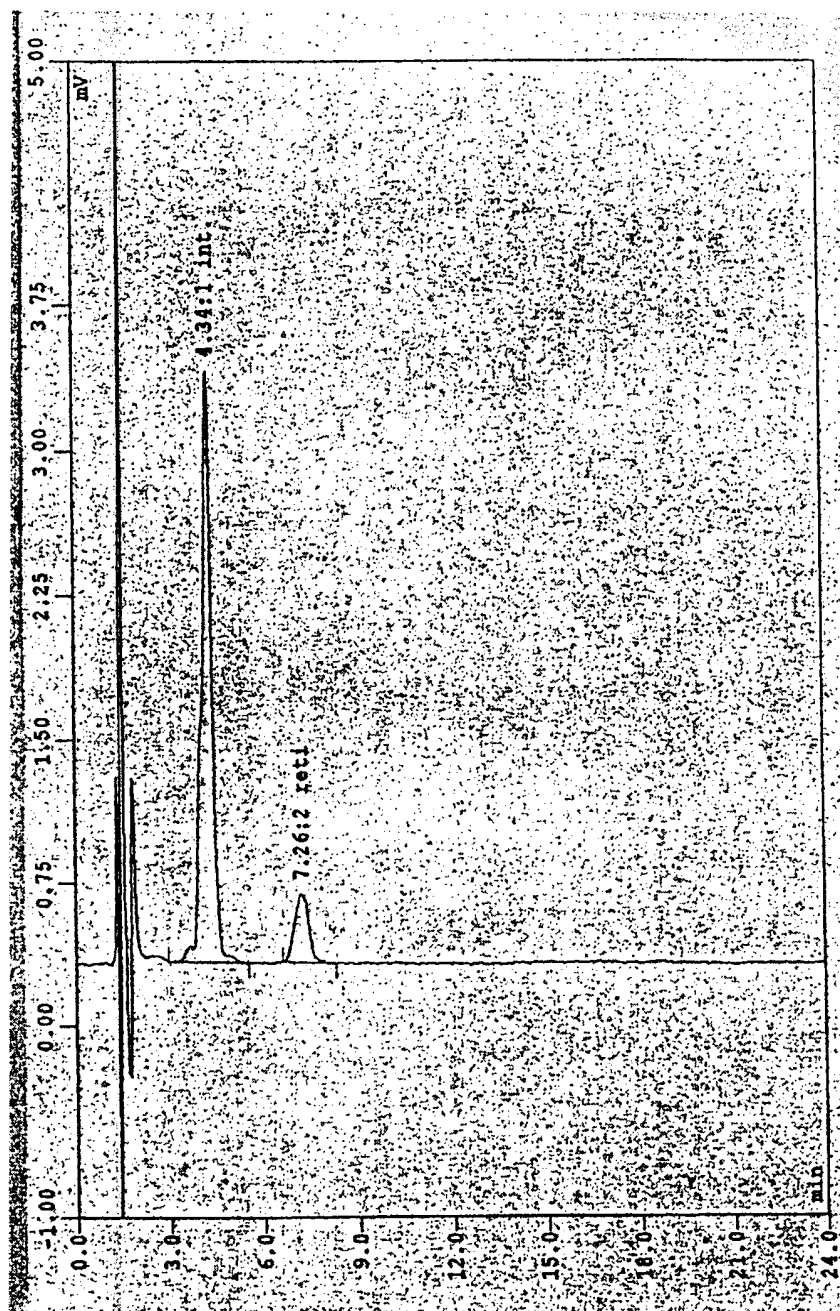


Fig. 7 shows an HPLC profile of the reaction mixture at the end of an activity assay for the β,β -carotene 15,15'-monooxygenase following the procedure in example 1. The first peak in the chromatogram represents the internal standard, while the second peak corresponds to retinal as the only product formed during the central cleavage with β -carotene as substrate.

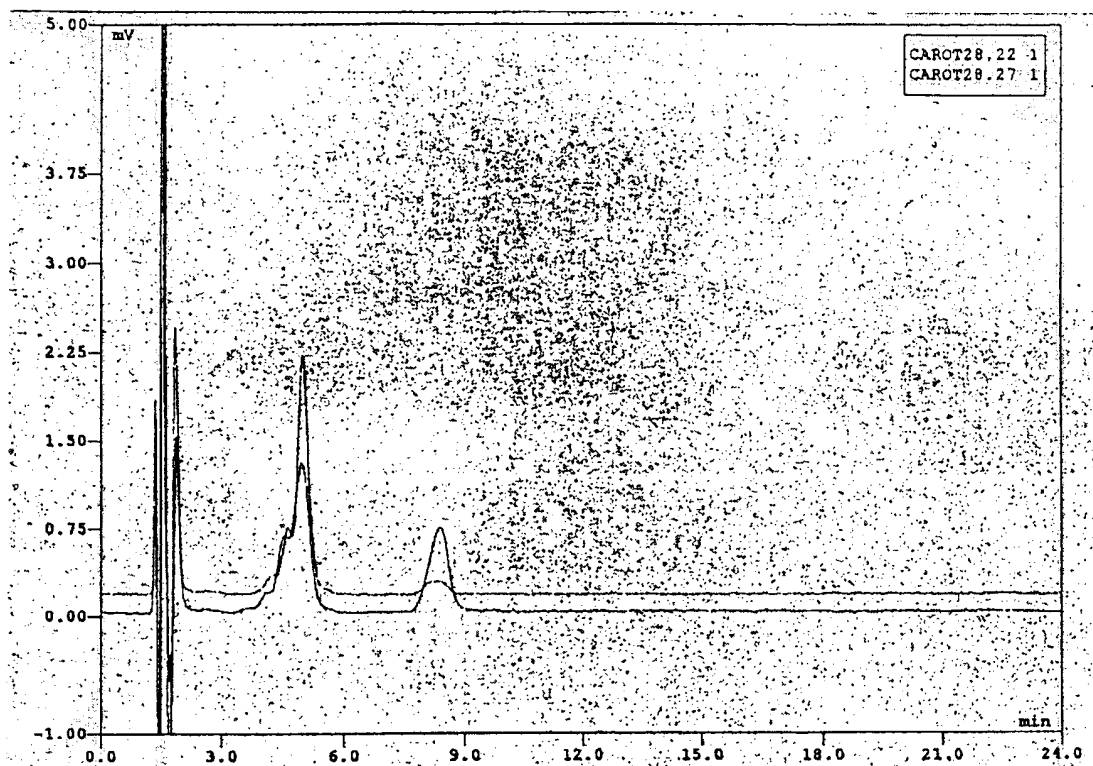


Fig. 8 confirms that the product peak in Fig. 7 is indeed retinal. A sample which was positive in the activity assay - (green (upper) chromatogram) was spiked with retinal and analysed in second HPLC run (red (lower) chromatogram). The chromatograms of the two runs were then overlayed.